

Fig. 1

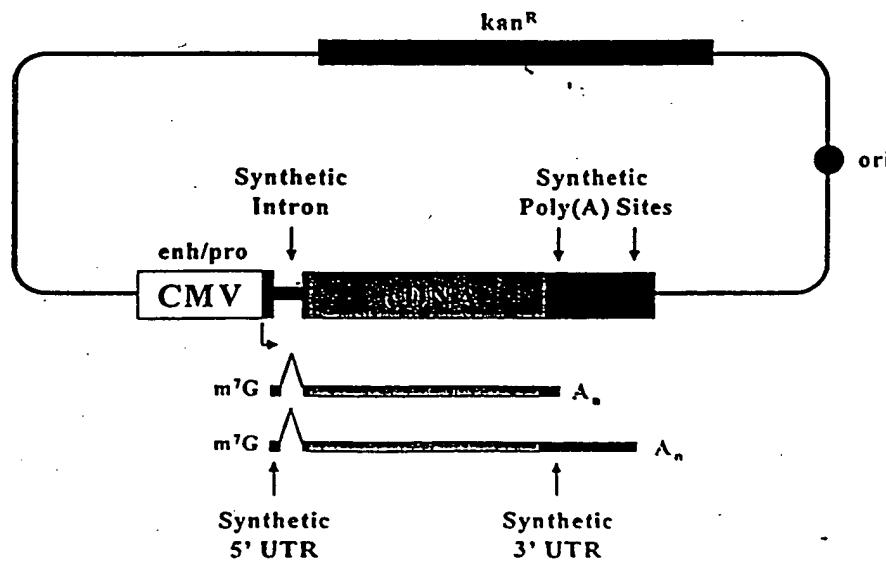


Fig. 2

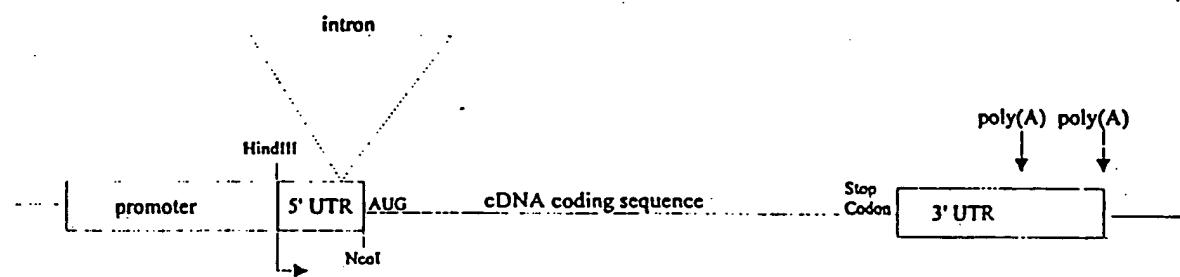


Fig. 3

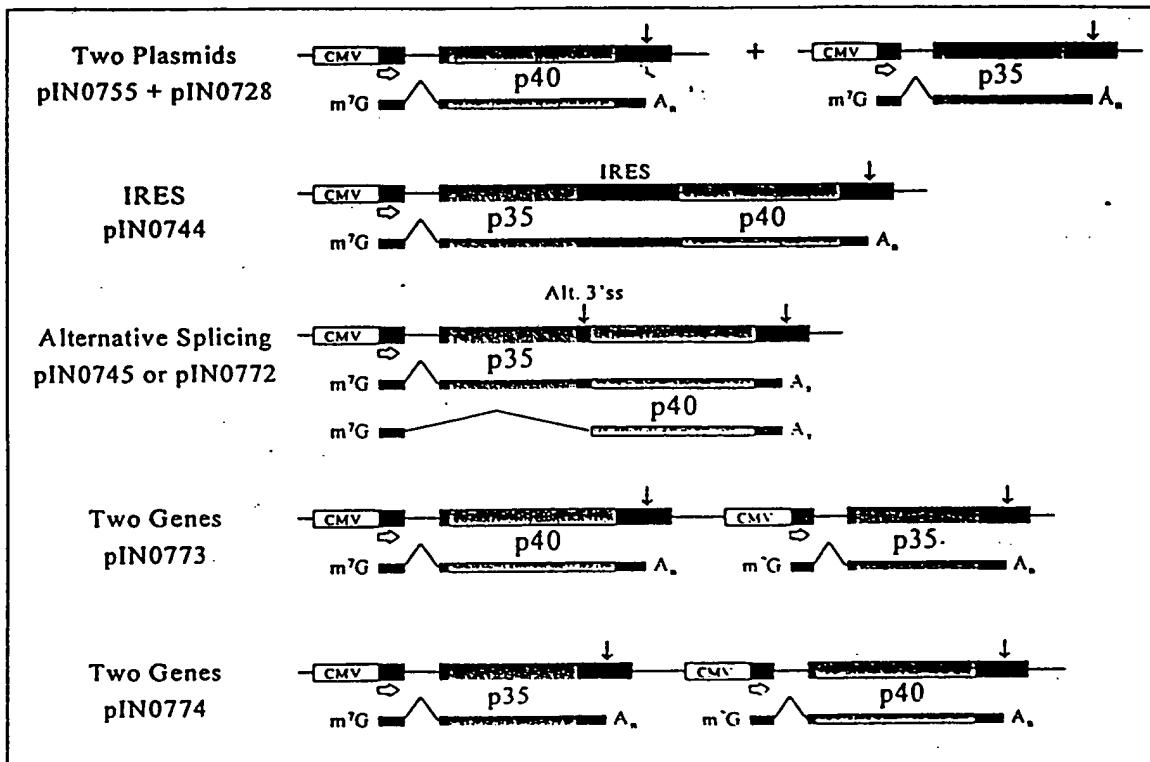


Fig. 4A

File : H40.AMI
 Range : 1 329
 Codon Table : Universal

SEQ ID NO. 1

10 20
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 ATG TGY CAY CAR CAR YTN GTN ATH WSN TGG TTY WSN YTN GTN TTY YTN GCN WSN CCN YTN

ATG TGT CAT CAA CAA TTA GTT ATT TCT TGG TTT TCT TTA GTT TTT TTA GCT TCT CCT TTA
 TGC CAC CAG CAG TTG GTC ATC TCC TTC TCC TTG GTC TTC TTG GCC TCC CCC TTG
 CTT GTA ATA TCA TCA CTT GTA CTT GCA TCA CCA CTT
 CTC GTG TCG TCG CTC GTG CTC GCG TCG CCG CTC
 CTA AGT AGT CTA CTA AGT CTA CTA
 CTG AGC AGC CTG AGC AGC CTG

30 40
 Val Ala Ile Trp Glu Leu Lys Lys Asp Val Tyr Val Val Glu Leu Asp Trp Tyr Pro Asp
 GTN GCN ATH TGG GAR YTN AAR AAR GAY GTN TAY GTN GTN GAR YTN GAY TGG TAY CCN GAY

GTG GCT ATT TGG GAA TTA AAA AAA GAT GTT TAT GTT GTT GAA TTA GAT TGG TAT CCT GAT
 GTC GCC ATC GAG TTG AAG AAG GAC GTC TAC GTC GTC GAG TTG GAC TAC CCC GAC
 GTA GCA ATA CTT GTA GTA CTT CCA
 GTG GCG CTC GTG GTG CTC CCG
 CTA CTG CTA CTG

50 60
 Ala Pro Gly Glu Met Val Val Leu Thr Cys Asp Thr Pro Glu Glu Asp Gly Ile Thr Trp
 GCN CCN GGN GAR ATG GTN YTN ACN TGY GAY ACN CCN GAR GAR GAY GGN ATH ACN TGG

GCT CCT GGT GAA ATG GTT TTA ACT TGT GAT ACT CCT GAA GAA GAT GGT ATT ACT TGG
 GCC CCC GGC GAG GTC GTC TTG ACC TGC GAC ACC CCC GAG GAG GAC GGC ATC ACC
 GCA CCA GGA GTA GTA CTT ACA ACA CCA GGA ATA ACA
 GCG CCG GGG GTG GTG CTC ACG ACG CCG GGG ACAC GGG
 CTA CTG

70 80
 Thr Leu Asp Gln Ser Ser Glu Val Leu Gly Ser Gly Lys Thr Leu Thr Ile Gln Val Lys
 ACN YTN GAY CAR WSN WSN GAR GTN YTN GGN WSN GGN AAR ACN YTN ACN ATH CAR GTN AAR

ACT TTA GAT CAA TCT TCT GAA GTT TTA GGT TCT GGT AAA ACT TTA ACT ATT CAA GTT AAA
 ACC TTG GAC CAG TCC TCC GAG GTC TTG GGC TCC GGC AAG ACC TTG ACC ATC CAG GTC AAG
 ACA CTT TCA TCA GTA CTT GGA TCA GGA ACA CTT ACA ATA GTA
 ACG CTC TCG TCG GTG CTC GGG TCG GGG ACG CTC ACG GTG
 CTA AGT AGT CTA AGT CTA
 CTG AGC AGC CTG AGC CTG

90 100
 Glu Phe Gly Asp Ala Gly Gln Tyr Thr Cys His Lys Gly Gly Glu Val Leu Ser His Ser
 GAR TTY GGN GAY GCN GGN CAR TAY ACN TGY CAY AAR GGN GGN GAR GTN YTN WSN CAY WSN

GAA TTT GGT GAT GCT GGT CAA TAT ACT TGT CAT AAA GGT GGT GAA GTT TTA TCT CAT TCT
 GAG TTC GGC GAC GCC GGC CAG TAC ACC TGC CAC AAG GGC GGC GAG GTC TTG TCC CAC TCC
 GGA GCA GGA ACA BGA BGA GTA CTT TCA TCA
 GGG GCG GGG ACG BGG BGG GTG CTC TCG TCG
 CTA AGT AGT CTA AGT
 CTG AGC AGC CTG AGC

Fig. 4B

110 120

Leu Leu Leu Leu His Lys Lys Glu Asp Gly Ile Trp Ser Thr Asp Ile Leu Lys Asp Gln
 YTN YTN YTN YTN CAY AAR AAR GAR GAY GGN ATH TGG WSN ACN GAY ATH YTN AAR GAY CAR

TTA TTA TTA TTA CAT AAA AAA GAA GAT GGT ATT TGG TCT ACT GAT ATT TTA AAA GAT CAA
 TTG TTG TTG TTG CAC AAG AAG GAG GAC GGC ATC TCC ACC GAC ATC TTG AAG GAC CAG
 CTT CTT CTT CTT GGA ATA TCA ACA ATA CTT
 CTC CTC CTC CTC GGG TCG ACG CTC
 CTA CTA CTA CTA AGT CTA
 CTG CTG CTG CTG AGC CTG

130 140

Lys Glu Pro Lys Asn Lys Thr Phe Leu Arg Cys Glu Ala Lys Asn Tyr Ser Gly Arg Phe
 AAR GAR CCN AAR AAY AAR ACN TTY YTN MGN TGY GAR GCN AAR AAY TAY WSN GGN MGN TTY

AAA GAA CCT AAA AAT AAA ACT TTT TTA CGT TGT GAA GCT AAA AAT TAT TCT GGT CGT TTT
 AAG GAG CCC AAG AAC AAG ACC TTC TTG CGC TGC GAG GCC AAG AAC TAC TCC GCC CGC TTC
 CCA ACA CTT CGA GCA TCA GGA CGA
 CCG ACG CTC CGG GCG TCG GGG CGG
 CTA AGA AGT AGA
 CTG AGG AGC AGG

150 160

Thr Cys Trp Trp Leu (Thr Thr) Ile Ser Thr Asp Leu Thr Phe Ser Val Lys Ser Ser Arg
 ACN TGY TGG TGG YTN ACN ACN ATH WSN ACN GAY YTN ACN TTY WSN GTN AAR WSN WSN MGN

ACT TGT TGG TGG TTA ACT ACT ATT TCT ACT GAT TTA ACT TTT TCT GTT AAA TCT TCT CGT
 ACC TGC TTG ACC ACC ATC TCC ACC GAC TTG ACC TTC TCC GTC AAG TCC TCC CGC
 ACA CTT ACA ACA ATA TCA ACA CTT ACA TCA GTA TCA TCA CGA
 ACG CTC ACG ACG TCG ACG CTC ACG TCG GTG TCG TCG CGG
 CTA AGT CTA AGT AGT AGT AGT AGA
 CTG AGC CTG AGC AGC AGC AGC AGG

170 180

Gly Ser Ser Asp Pro Gln Gly Val Thr Cys Gly Ala Ala Thr Leu Ser Ala Glu Arg Val
 GGN WSN WSN GAY CCN CAR GGN GTN ACN TGY GGN GCA GCN ACN YTN WSN GCN GAR MGN GTN

GGT TCT TCT GAT CCT CAA GGT GTT ACT TGT GGT GCT GCT ACT TTA TCT GCT GAA CGT GTT
 GGC TCC TCC GAC CCC CAG GGC GTC ACC TGC GGC GCC ACC TTG TCC GCC GAG CGC GTC
 GGA TCA TCA CCA GGA GTA ACA TGA GCA GCA ACA CTT TCA GCA CGA GTA
 GGG TCG TCG CCG GGG GTG ACG CGG GCG GCG ACG CTC TCG GCG CGG GTG
 AGT AGT CTA AGT AGA
 AGC AGC CTG AGC AGG

190 200

Arg Gly Asp Asn Lys Glu Tyr Glu Tyr Ser Val Glu Cys Gln Glu Asp Ser Ala Cys Pro
 MGN GGN GAY AAY AAR GAR TAY GAR TAY WSN GTN GAR TGY CAR GAR GAY WSN GCN TGY CCN

CGT GGT GAT AAT AAA GAA TAT GAA TAT TGT TTG GAA TGT CAA GAA GAT TCT GCT TGT CCT
 CGC GGC GAC AAC AAG GAG TAC AG TAC TGC GAT TGC CAG GAC TCC GCC TGC CCC
 CGA GGA TCA GTA TCA GCA CCA
 CGG GGG TCG GTG TCG GCG CCG
 AGA AGT AGT
 AGG AGC AGC AGC

Fig. 4C

210

Ala Ala Glu Glu Ser Leu Pro Ile Glu Val Met Val Asp Ala Val His Lys Leu Lys Tyr
GCU GCU GAR GAR WSN YTN CCN ATH GAR GTN ATG GTN GAY GCU GTN CAY AAR YTN AAR TAY

GCT GCT GAA GAA TCT TTA CCT ATT GAA GTT ATG GTT GAT GCT GTC CAT AAA TTA AAA TAT
GCC GCC GAG GAG TCC TTG CCC ATC GAG GTC GTC GAC GCC GTC CAC AAG TTG AAG TAC
GCA GCA TCA CTT CCA ATA GTA GTA GCA GTA CTT
GCG GCG TCG CTC CCG GTG GTG GCG GTG CTC
AGT CTA CTA
AGC CTG CTG

220

Glu Asn Tyr Thr Ser Ser Phe Phe Ile Arg Asp Ile Ile Lys Pro Asp Pro Pro Lys Asn
GAR AAY TAY ACN WSN WSN TTG TTG ATH MGN TAY ATH ATH AAR CCA GAY CCN CCN AAR AAY

GAA AAT TAT ACT TCT TCT TTT ATT CGT AT ATT ATT AAA CC GAT CCT CCT AAA AAT
GAG AAC TAC ACC TCC TCC TTC ATC CGC AAC ATC ATC ATC AAG CC GAC CCC CCC AAG AAC
ACA TCA TCA ATA CGA ATA ATA CGA CCA CCA
ACG TCG TCG CGA CGA CGA CCG CCG
AGT AGT AGA AGA
AGC AGC AGG AGG

230

Leu Gln Leu Lys Pro Leu Lys Asn Ser Arg Gln Val Glu Val Ser Trp Glu Tyr Pro Asp
YTN CAR YTN AAR CCN YTN AAR AAY WSN MGN CAR GTN GAR GTN WSN TGG GAR TAY CCN GAY

TTA CAA TTA AAA CCT TTA AAA AAT TCT CGT CAA GAA GTT TCT TGG GAA TAT CCT GAT
TTG CAG TTG AAG CCC TTG AAG AAC TCC CCC CAG GTC GAG GTC TCC GAG TAC CCC GAC
CTT CTT CCA CTT TCA CGA GTA GTA TCA CCA
CTC CTC CCG CTC TCG CGG GTG GTG TCG CCG
CTA CTA CTA AGT AGA AGT
CTG CTG CTG AGC AGG AGC

250

Leu Gln Leu Lys Asn Ser Arg Gln Val Glu Val Ser Trp Glu Tyr Pro Asp
YTN CAR YTN AAR CCN YTN AAR AAY WSN MGN CAR GTN GAR GTN WSN TGG GAR TAY CCN GAY

TTA CAA TTA AAA CCT TTA AAA AAT TCT CGT CAA GAA GTT TCT TGG GAA TAT CCT GAT
TTG CAG TTG AAG CCC TTG AAG AAC TCC CCC CAG GTC GAG GTC TCC GAG TAC CCC GAC
CTT CTT CCA CTT TCA CGA GTA GTA TCA CCA
CTC CTC CCG CTC TCG CGG GTG GTG TCG CCG
CTA CTA CTA AGT AGA AGT
CTG CTG CTG AGC AGG AGC

270

Thr Trp Ser Thr Pro His Ser Tyr Phe Ser Leu Thr Phe Cys Val Gln Val Gln Gly Lys
ACN TGG WSN ACN CCN CAY WSN TAY TTG WSN YTN ACN TTY TGY GTN CAR GTN CAR GGN AAR

ACT TGG TCT ACT CCT CAT TCT TAT TTT TCT TTA ACT TTT TGT GTT CAA GAA GTT CAA GGT AAA
ACC TCC ACC CCC CAC TCC TAC TTC TCC TTG ACC TTC TGC GTC CAG GTC CAG GGC AAG
ACA TCA ACA CCA TCA TCA CTT ACA GTA GTA GGA
ACG TCG ACG CCG TCG TCG CTC ACG GTG GTG GGG
AGT AGT AGT CTA
AGC AGC AGC AGC CTG

280

Thr Trp Ser Thr Pro His Ser Tyr Phe Ser Leu Thr Phe Cys Val Gln Val Gln Gly Lys
ACN TGG WSN ACN CCN CAY WSN TAY TTG WSN YTN ACN TTY TGY GTN CAR GTN CAR GGN AAR

ACT TGG TCT ACT CCT CAT TCT TAT TTT TCT TTA ACT TTT TGT GTT CAA GAA GTT CAA GGT AAA
ACC TCC ACC CCC CAC TCC TAC TTC TCC TTG ACC TTC TGC GTC CAG GTC CAG GGC AAG
ACA TCA ACA CCA TCA TCA CTT ACA GTA GTA GGA
ACG TCG ACG CCG TCG TCG CTC ACG GTG GTG GGG
AGT AGT AGT CTA
AGC AGC AGC AGC CTG

290

Ser Lys Arg Glu Lys Lys Asp Arg Val Phe Thr Asp Lys Thr Ser Ala Thr Val Ile Cys
WSN AAR MGN GAR AAR AAR GAY MGN GTN TTY ACN GAY AAR ACN WSN GCU ACN GTN ATH TGY

TCT AAA CGT GAA AAA AAA GAT CGT GTT TTT ACT GAT AAA ACT TCT GCT ACT GTT ATT TGT
TCC AAG CGC GAG AAG AAG GAC CGC GTC TTC ACC GAC AAG ACC TCC GCC ACC GTC ATC TGC
TCA CGA CGA GTA ACA ACA TCA GCA ACA GTA ATA
TCG CGG CGG GTG ACG ACG TCG GCG ACG GTG
AGT AGA AGA AGT
AGC AGG AGG AGC

300

Fig. 4D

310

Arg Lys Asn Ala Ser Ile Ser Val Arg Ala Gln Asp Arg Tyr Tyr Ser Ser Ser Trp Ser
 MGN AAR AAY GCN WSN ATH WSN GTN MGN GCN CAR GAY MGN TAY TAY WSN WSN WSN TGG WSN

CGT	AAA	AAT	GCT	TCT	ATT	TCT	GTT	CGT	GCT	CAA	GAT	CGT	TAT	TAT	TCT	TCT	TCT	TGG	TCT
CGC	AAG	AAC	GCC	TCC	ATC	TCC	GTC	CGC	GCC	QAG	GAC	CGC	TAC	TAC	TCC	TCC	TCC		TCC
CGA		GCA	TCA	ATA	TCA	GTA	CGA	GCA			CGA		TCA	TCA	TCA			TCA	
CGG		GCG	TCG		TCG	GTG	CGG	GCG			CGG		TCG	TCG	TCG			TCG	
AGA			AGT		AGT		AGA			AGA		AGT	AGT	AGT				AGT	
AGG			AGC		AGC		AGG			AGG		AGC	AGC	AGC				AGC	

Glu Trp Ala Ser Val Pro Cys Ser ***
GAR TGG GCG WSN GTN CCN TGY WSN TRR

GAA	TGG	GCT	TCT	GTT	CCT	TGT	TCT	TAA
GAG		GCC	TCC	GTC	CCC	TGG	TCC	TAG
		GCA	TCA	GTA	CCA		TCA	TGA
		GCG	TCG	GTG	CCG		TCG	
			AGT				AGT	
			AGC				AGC	

Fig. 5A

File : H35.AMI
 Range : 1 220
 Codon Table : Universal

SEQ 10 NO. 5

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Met Cys Pro Ala Arg Ser Leu Leu Leu Val Ala Thr Leu Val Leu Leu Asp His Leu Ser
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 ATG TGT CCT GCT CGT TCT TTA TTA GTT GCT ACT TTA GTT TTA TTA GAT CAT TTA TCT
 TGC CCC GCC CGC TCC TTG TTG GTC GCC ACC TTG GTC TTG TTG GAC CAC TTG TCC
 CCA GCA CGA TCA CTT CTT CTT GTC GCA ACA CTT GTC CTT CTT CTT TCA
 CCG GCG CGG TCG CTC CTC GTG GCG ACG CTC GTG CTC CTC CTC TCG
 AGA AGT CTA CTA CTA CTA CTA CTA CTA CTA CTA AGT CTA AGT
 AGG AGC CTG CTG CTG CTG CTG CTG CTG AGC

30

40

Leu Ala Arg Asn Leu Pro Val Ala Thr Pro Asp Pro Gly Met Phe Pro Cys Leu His His
 YTN GCN MGN AAY YTN CCN GTN GCN ACN CCN GAY CCN GGN ATG TTY CCN TGY YTN CAY CAY

 TTA GCT CGT AAT TTA CCT GTT GCT ACT CCT GAT CCT GGT ATG TTT CCT TGT TTA CAT CAT
 TTG GCC CGG AAC TTG CCC GTC GCC ACC CCC GAC CCC GGC TTC CCC TGC TTG CAC CAC
 CTT GCA CGA CTT CCA GTC GCA ACA CCA CCA GGA CCA CTT
 CTC CGG CGG CTC CCG GTG GCG ACG CCG CCG GGG CCG CTC CTA
 CTA AGA CTA
 CTG AGG CTG CTG CTG CTG CTG CTG CTG CTG

50

60

Ser Gln Asn Leu Leu Arg Ala Val Ser Asn Met Leu Gln Lys Ala Arg Gln Thr Leu Glu
 WSN CAR AAY YTN YTN MGN GCN GTN WSN AAY ATG YTN CAR AAR GCN MGN CAR ACN YTN GAR

 TCT CAA AAT TTA TTA CGT GCT GTT TCT AAT ATG TTA CAA AAA GCT CGT CAA ACT TTA GAA
 TCC CAG AAC TTG TTG CGC GCC GTC TCC AAC TTG CAG AAG GCC CGC CAG ACC TTG GAG
 TCA CTT CTT CGA GCA GTC TCA CTT GCA CCA ACA CTT
 TCG CTC CTC CGG GCG GTG TCG CTC GCG CGG ACA CTT
 AGT CTA CTA AGA AGT CTA AGA CTA AGC CTA AGA CTA
 AGC CTG CTG AGG AGC CTG AGG AGC CTG AGC CTG

70

80

Phe Tyr Pro Cys Thr Ser Glu Glu Ile Asp His Glu Asp Ile Thr Lys Asp Lys Thr Ser
 TTY TAY CCN TGY ACN WSN GAR GAR ATH GAY CAY GAR GAY ATH ACN AAR GAY AAR ACN WSN

 TTT TAT CCT TGT ACT TCT GAA GAA ATT GAT CAT GAA GAT ATT ACT AAA GAT AAA ACT TCT
 TTC TAC CCC TGC ACC TCC GAG GAG ATC GAC CAC GAG GAC ATC ACC AAG GAC AAG ACC TCC
 CCA ACA TCA ATA ATA ACA .. ACA TCA
 CCG ACG TCG AGT ACG ACG TCG AGT
 AGC AGC AGC AGC

90

100

Thr Val Glu Ala Cys Leu Pro Leu Glu Leu Thr Lys Asn Glu Ser Cys Leu Asn Ser Arg
 ACN GTN GAR GCN TGY YTN CCN YTN GAR YTN ACN AAR AAY GAR WSN TGY YTN AAY WSN MGN

 ACT GTT GAA GCT TGT TTA CCT TTA GAA TTA ACT AAA ATT GAA TCT TGT TTA AAT TCT CGT
 ACC GTC GAG GCC TGC TTG CCC TTG GAG TTG ACC AAG AAC GAG TCC TGC TTG AAC TCC CGC
 ACA GTA GCA CTT CCA CTT CTT ATA TCA CTT CTC CGA
 ACG GTG GCG CTC CCC TC CTC ACG 3 CTC TCG CGG
 CTA CTA CTA AGT CTA AGT
 CTG CTG CTG AGC CTG AGC AGC AGG

Fig. 5B

110

120

Glu Thr Ser Phe Ile Thr Asn Gly Ser Cys Leu Ala Ser Arg Lys Thr Ser Phe Met Met
 GAR ACN WSN TTY ATH ACN AAY GGN WSN TGY YTN GCN WSN MGN AAR ACN WSN TTY ATG ATG

GAA ACT TCT TTT ATT ACT AAT GGT TCT TGT TTA GCT TCT CGT AAA ACT TCT TTT ATG ATG
 GAG ACC TCC TTC ATC ACC AAC GGC TCC TGC TTG GCC TCC CGC AAG ACC TCC TTC
 ACA TCA ATA ACA GGA TCA CTT GCA TCA CGA ACA TCA
 ACG TCG ACG GGG TCG CTC GCG TCG CGG ACG TCG
 AGT AGT CTA AGT AGA AGT
 AGC AGC CTG AGC AGG AGC

130

140

Ala Leu Cys Leu Ser Ser Ile Tyr Glu Asp Leu Lys Met Tyr Glu Val Glu Phe Lys Thr
 GCN YTN TGY YTN WSN WSN ATH TAY GAR GAY YTN AAR ATG TAY CAR GTN GAR TTY AAR ACN

GCT TTA TGT TTA TCT TCT ATT TAT GAA GAT TTA AAA ATG TAT CAA GTT GAA TTT AAA ACT
 GCC TTG TGC TTG TCC TCC ATC TAC GAG GAC TTG AAG TAC CAG GTC GAG TTC AAG ACC
 GCA CTT CTT TCA TCA ATA CTT GTA ACA
 GCG CTC CTC TCG TCG CTC GTG ACG
 CTA CTA AGT AGT CTA
 CTG CTG AGC AGC CTG

150

160

Met Asn Ala Lys Leu Leu Met Asp Pro Lys Arg Gln Ile Phe Leu Asp Gln Asn Met Leu
 ATG AAY GCN AAR YTN YTN ATG GAY CCN AAR MGN CAR ATH TTY YTN GAY CAR AAY ATG YTN

ATG AAT GCT AAA TTA TTA ATG GAT CCT AAA CGT CAA ATT TTT TTA GAT CAA AAT ATG TTA
 AAC GCC AAG TTG TTG GAC CCC AAG CGC CAG ATC TTC TTG GAC CAG AAC TTG
 GCA CTT CTT CCA CGA ATA CTT CTT
 GCG CTC CTC CCG CGG CTC CTC
 CTA CTA AGA CTA CTA
 CTG CTG AGG CTG CTG

170

180

Ala Val Ile Asp Glu Leu Met Gln Ala Leu Asn Phe Asn Ser Glu Thr Val Pro Gln Lys
 GCN GTN ATH GAY GAR YTN ATG CAR GCN YTN AAY TTY AAY WSN GAR ACN GTN CCN CAR AAR

GCT GTT ATT GAT GAA TTA ATG CAA GCT TTA AAT TTT AAT TCT GAA ACT GTT CCT CAA AAA
 GCC GTC ATC GAC GAG TTG CAG GCC TTG AAC TTC AAC TCC GAG ACC GTC CCC CAG AAG
 GCA GTA ATA CTT GCA CTT TCA ACA GTA CCA
 GCG GTG CTC GCG CTC TCG ACG GTG CCG
 CTA CTA AGT
 CTG CTG AGC

190

200

Ser Ser Leu Glu Glu Pro Asp Phe Tyr Lys Thr Lys Ile Lys Leu Cys Ile Leu His
 WSN WSN YTN GAR GAR CCN GAY TTY TAY AAR ACN AAR ATH AAR YTN TGY ATH YTN YTN CAY

TCT TCT TTA GAA GAA CCT GAT TTT TAT ATA ACT AAA ATT AAA TTA TGT ATT TTA TTA CAT
 TCC TCC TTG GAG GAG CCC GAC TC TAC AAG ACC AAG ATC AAG 3 TGC ATC TTG TTG CAC
 TCA TCA CTT CCA ACA ATA CTT ATA CTT CTT
 TCG TCG CTC CCG ACG CTC CTC CTC
 AGT AGT CTA CTA CTA CTA
 AGC AGC CTG CTG CTG CTG

Fig. 5C

210

220

Ala Phe Arg Ile Arg Ala Val Thr Ile Asp Arg Val Thr Ser Tyr Leu Asn Ala Ser ***
GCN TTY MGN ATH MGN GCN GTN ACN ATH GAY MGN GTN ACN WSN TAY YTN AAY GCN WSN TRR

GCT TTT CGT ATT CGT GCT GTT ACT ATT GAT CGT GTT ACT TCT TAT TTA AAT GCT TCT TAA
GCC TTC CGC ATC CGC GCC GTC ACC ATC GAC CGC GTC ACC TCC TAC TTG AAC GCC TCC TAG
GCA CGA ATA CGA GCA GTA ACA ATA CGA GTA ACA TCA CTT GCA TCA TGA
GCG CGG CGG GCG GTG ACG CGG GTG ACG TCG CTC GCG TCG
AGA AGA AGA AGT CTA AGT
AGG AGG AGG AGC CTG AGC

Fig. 6

Codon Frequency Tables

human high.cod

Codon usage for human (highly expressed) genes 1/24/91.

AAcid	Codon	Number	/1000	Fraction
Gly	GCG	965.00	18.76	0.24
Gly	GCA	525.00	10.18	0.14
Gly	GCT	441.00	9.14	0.12
Gly	GCC	1867.00	38.70	0.50
Cle	GAA	2426.00	50.18	0.73
Glu	CAA	732.00	14.62	0.25
Asp	CAT	532.00	12.27	0.25
Asp	CAC	1021.00	21.73	0.75
Val	GTC	1866.00	38.68	0.64
Val	GTA	134.00	2.78	0.05
Val	GTT	191.00	4.10	0.07
Val	GTC	721.00	15.49	0.25
Ala	GCG	652.00	13.51	0.17
Ala	GCA	486.00	10.12	0.13
Ala	GCT	634.00	13.56	0.17
Ala	GCC	2651.00	42.64	0.53
Arg	AGG	512.00	10.61	0.18
Arg	AGA	294.00	6.18	0.10
Ser	ACT	354.00	7.34	0.10
Ser	ACC	1171.00	24.27	0.36
Lys	AAA	2117.00	43.28	0.82
Lys	AAU	471.00	9.76	0.18
Lys	AAT	314.00	6.51	0.22
Lys	AAC	1120.00	23.22	0.78
Met	ATG	1077.00	22.32	1.00
Ile	ATA	88.00	1.82	0.05
Ile	ATT	315.00	6.53	0.21
Ile	ATC	1369.00	28.38	0.77
Thr	ACG	465.00	9.10	0.15
Thr	ACA	273.00	5.73	0.14
Thr	ACT	358.00	7.62	0.14
Thr	ACC	1502.00	31.13	0.57
Ter	TCC	652.00	13.51	1.00
Ter	TCA	109.00	2.26	0.05
Cys	TCT	325.00	6.74	0.32
Cys	TCC	706.00	14.63	0.64
End	TAC	42.00	0.87	0.21
End	TAA	46.00	0.95	0.23
Tyr	TAT	360.00	7.46	0.26
Tyr	TAC	1042.00	21.60	0.74
Lys	TTC	313.00	6.45	0.06
Lys	TTA	76.00	1.58	0.02
Pro	TTT	336.00	6.96	0.20
Pro	TTC	1377.00	28.54	0.80
Ser	TCC	329.00	6.74	0.09
Ser	TCA	165.00	3.42	0.05
Ser	TCT	456.00	9.33	0.13
Ser	TCC	958.00	19.86	0.28
Arg	CCC	611.00	12.67	0.21
Arg	CCA	183.00	3.79	0.06
Arg	CCT	216.00	4.35	0.07
Arg	CCG	1086.00	22.51	0.37
Cln	CAC	2020.00	41.17	0.68
Cln	CAA	283.00	5.17	0.12
His	CAT	234.00	4.15	0.21
His	CAC	870.00	17.03	0.39
Lys	CTG	2886.00	51.78	0.64
Lys	CTA	166.00	3.14	0.03
Lys	CTT	238.00	4.33	0.05
Lys	CTC	1276.00	26.45	0.26
Pro	CCC	162.00	3.09	0.17
Pro	CCA	456.00	9.43	0.26
Pro	CCT	568.00	11.77	0.22
Pro	CCC	1410.00	29.23	0.44

Fig. 7

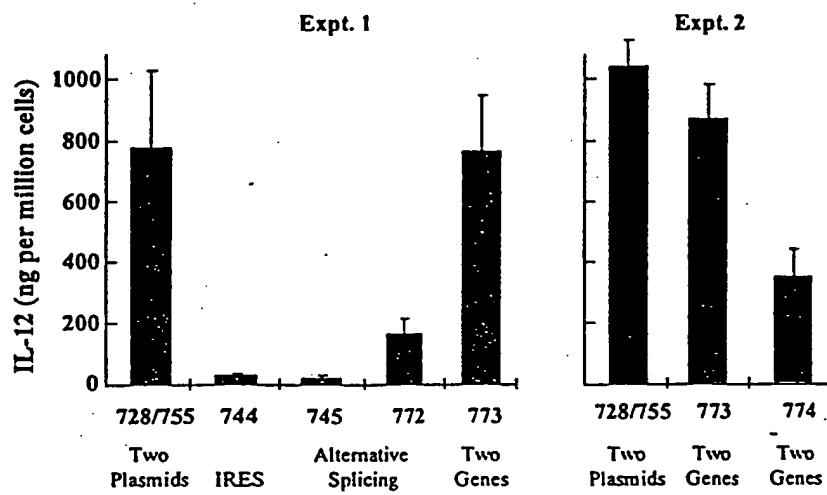


Fig. 8

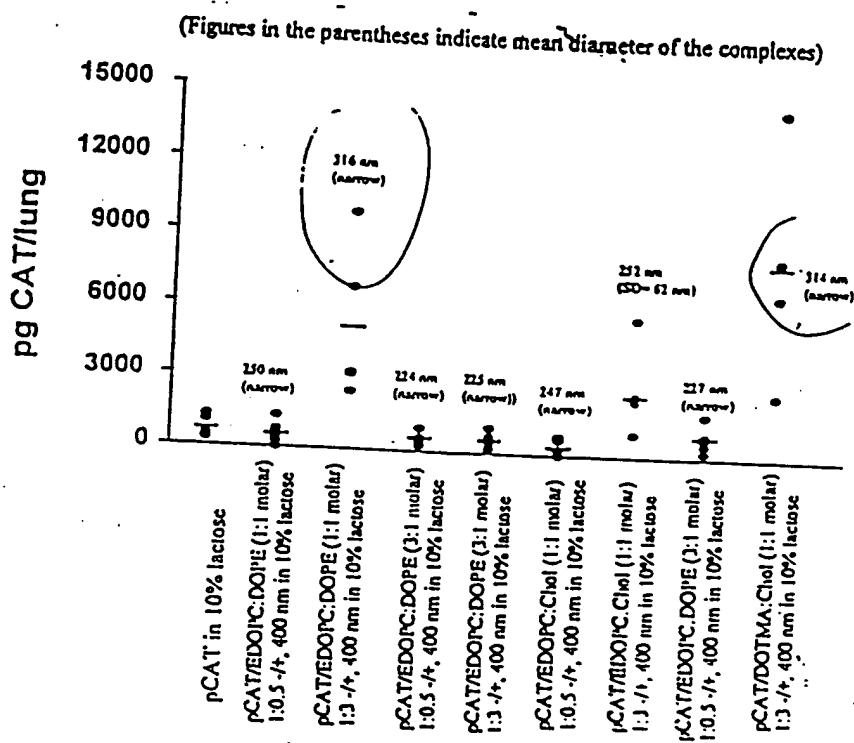
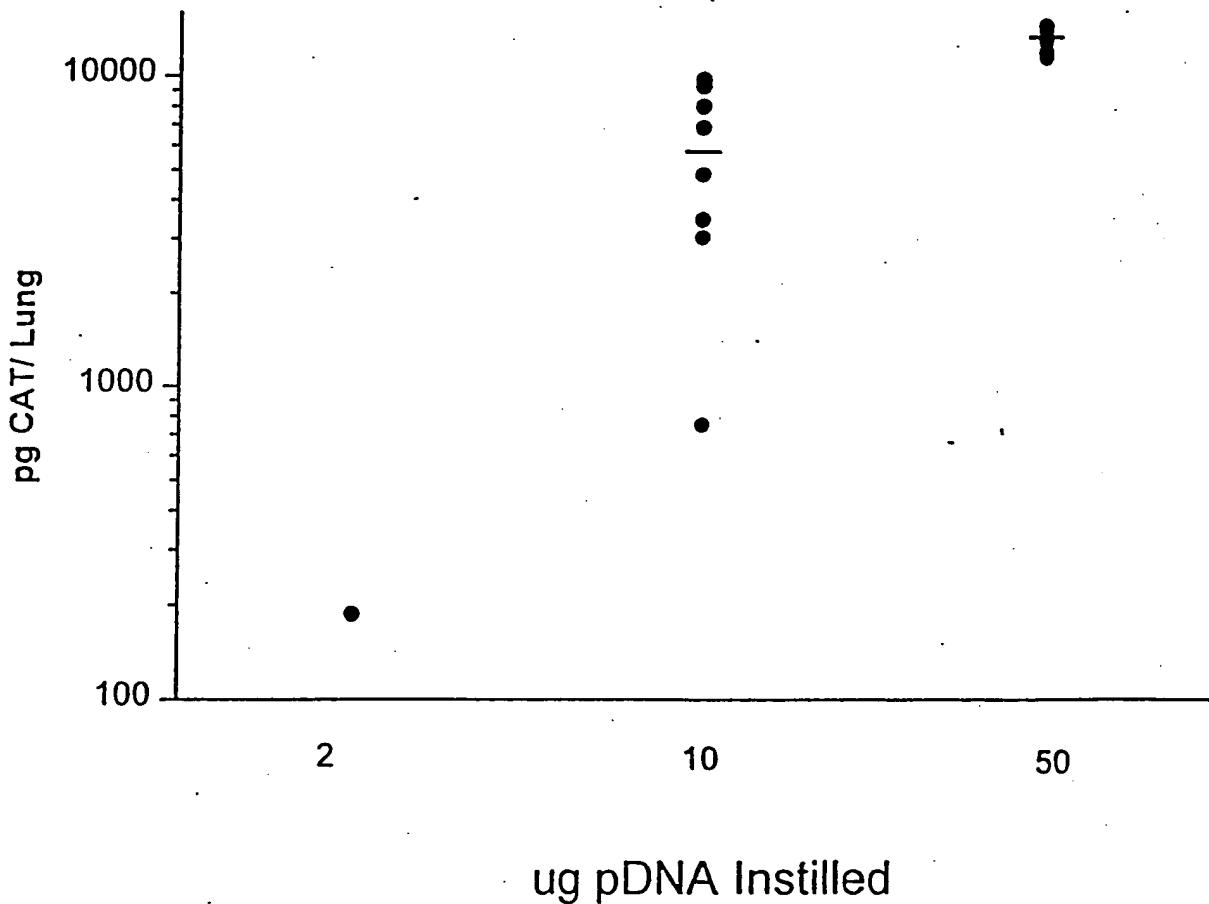


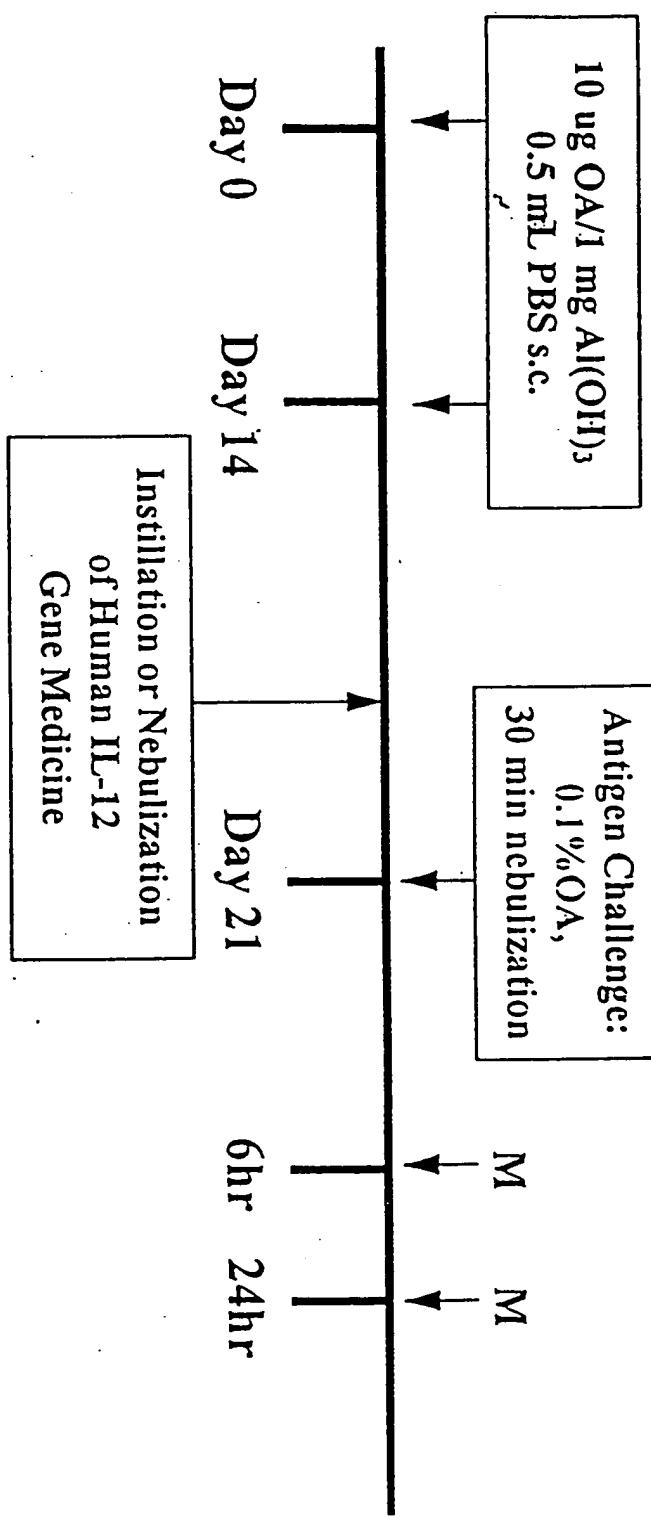
Fig. 9

(pCT0129.095:DOTMA/CHOL 1:3 -/+ 10% Lactose)



Antigen-Induced Airway Inflammation Model in Guinea Pigs

Fig. 10



M (measurement) = bronchoalveolar lavage total and differential cell count

Fig. 11

